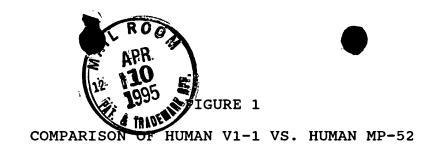
U 8/217780



V1-1 1 MP52 1	AGC GCT	CGC	TGC	AGC	CGC CGG	AAG AAG	CCG GCA	TTG CTG	CAC	GTG GTC	GAC AAC	TTC	AAG AAG	GAG GAC	CTC ATG
16 46 46 16	GGC	TGG TGG	GAC GAC	Asp GAC GAC Asp	TGG TGG	ATC ATC	ATC ATC	GCG GCA	CCG	CTG CTT	GAC GAG	TAC	GAG GAG	GCG GCT	TAC TTC
31 91 91 31	CAC	TGC	GAG GAG	Gly GGC GGG Gly	CTT CTG	TGC	GAC GAG	TTC	CCT	TTG	CGT CGC	TCG TCC	CAC CAC	CTC	GAG GAG
46 121 121 46	CCC	ACC ACG	AAC AAT	His CAT !!! CAT His	GCC GCA	ATC GTC	ATT ATC	CAG CAG	ACG	CTG CTG	CTC ATG	AAC AAC	TCC	ATG ATG	GCA GAC
61 181 181 61	CCA CCC	GAC GAG	GCG - TCC	Ala GCG - ACA Thr	CCG CCA	CCC	TCC ACC	TGC TGC	TGT TGT	GTG GTG	CCA CCC	GCG ACG	CGC CGG	CTC CTG	AGC AGT
76 226 226 76	CCC	ATC ATC	AGC AGC	Ile ATC ATC Ile	CTC	TAC TTC	ATC ATT	GAC GAC	GCC - TCT	GCC GCC	AAC AAC	AAC AAC	GTT GTG	GTC GTG	TAC
91 271 271 91	AAG AAG	CAA CAG	TAC	Glu GAG GAG Glu	GAC GAC	ATG ATG	GTC GTC	GTG GTG	GAG GAG	GCC - TCG	TGC TGT	GGC GGC	TGC TGC	AGG AGG	

Homology at the nucleotide level: 249/312 = 79.8%Homology at the amino acid level: 84/104 = 80.8%